<!--StartFragment-->RESULT 8
US-09-878-722-245
; Sequence 245, Application US/09878722
; Publication No. US20020040127A1
; GENERAL INFORMATION:
; APPLICANT: Jiang, Yuqiu
; APPLICANT: Hepler, William T.
; APPLICANT: Clapper, Jonathan
; APPLICANT: Wang, Aijun
; APPLICANT: Secrist, Heather
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
; TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
; FILE REFERENCE: 210121.524
; CURRENT APPLICATION NUMBER: US/09/878,722
; CURRENT FILING DATE: 2001-06-08
; NUMBER OF SEQ ID NOS: 245
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 245
;   LENGTH: 564
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-878-722-245

   Query Match   98.7%; Score 2716.5; DB 3; Length 564;
   Best Local Similarity   91.1%; Pred. No. 3.5e-271;
   Matches 514; Conservative   1; Mismatches   0; Indels 49; Gaps   1;

Qy   1 MGNWVVNHWFSVLFLVVWLGLNVFLFVDAFLKYEKADKYYYTRKILGSTLACARASALCL 60
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1 MGNWVVNHWFSVLFLVVWLGLNVFLFVDAFLKYEKADKYYYTRKILGSTLACARASALCL 60

Qy   61 NFNSTLILLPVCRNLLSFLRGTCSFCSRTLRKQLDHNLTFHKLVAYMICLHTAIHIIAHL 120
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy   121 FNFDCYSRSRQATDGSLASILSSLSHDEKKGGSWLNPIQSRNTTVEYVTFTSIAGLTGVI 180
            ||||||||||||||||||||||||||||||||||||||||||||||||||:||||||
Db   121 FNFDCYSRSRQATDGSLASILSSLSHDEKKGGSWLNPIQSRNTTVEYVTFTSVAGLTGVI 180

Qy   181 MTIALILMVTSATEFIRRSYFEVFWYTHHLFIFYILGLGIHGIGGIVRGQTEESMNESHP 240
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   181 MTIALILMVTSATEFIRRSYFEVFWYTHHLFIFYILGLGIHGIGGIVRGQTEESMNESHP 240

Qy   241 RKCAESFEMWDDRDSHCRRPKFEGHPPESWKWILAPVILYICERILRFYRSQQKVVITKV 300
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   241 RKCAESFEMWDDRDSHCRRPKFEGHPPESWKWILAPVILYICERILRFYRSQQKVVITKV 300

Qy   301 VMHPSKVLELQMNKRGFSMEVGQYIFVNCPSISLLEWHPFTLTSAPEEDFFSIHIRAAGD 360
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   301 VMHPSKVLELQMNKRGFSMEVGQYIFVNCPSISLLEWHPFTLTSAPEEDFFSIHIRAAGD 360

Qy   361 WTENLIRAFEQQYSPIPRIEVDGPFGTASEDVFQYEVAVLVGAGIGVTPFASILKSIWYK 420
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   361 WTENLIRAFEQQYSPIPRIEVDGPFGTASEDVFQYEVAVLVGAGIGVTPFASILKSIWYK 420

Qy   421 FQCADHNLKTKK--- 432
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Db   421 FQCADHNLKTKKIYFYWICRETGAFSWFNNLLTSLEQEMEELGKVGFLNYRLFLTGWDSN 480

Qy   433 -VGHAALNFDKATDIVTGLKQKTSFGRPMWDNEFSTIATSHPKSVVGVFLCGPRTLAKSL 491
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Qy      492 RKCCHRYSSLDPRKVQFYFNKENF 515
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Db      541 RKCCHRYSSLDPRKVQFYFNKENF 564
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